

100-443886-100

<223> Wherein Trp is D-tryptophan

<212> PRT
<213> *Cavia porcellus*

<400> 7
Arg Pro Lys Pro Gln Gln Phe Phe Gly Leu Met
1 5 10

<210> 8
<211> 11
<212> PRT
<213> *Gallus gallus*

<400> 8
Arg Pro Arg Pro Gln Gln Phe Phe Gly Leu Met
1 5 10

<210> 9
<211> 11
<212> PRT
<213> *Gadus morhua*

<400> 9
Lys Pro Arg Pro Gln Gln Phe Ile Gly Leu Met
1 5 10

<210> 10
<211> 11
<212> PRT
<213> *Oncorhynchus mykiss*

<400> 10
Lys Pro Arg Pro His Gln Phe Phe Gly Leu Met
1 5 10

<210> 11
<211> 10
<212> PRT
<213> *Petromyzon marinus*

<400> 11
Ala Lys His Asp Lys Phe Tyr Gly Leu Met
1 5 10

<210> 12
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<221> VARIANT
<222> (1)
<223> Wherein Xaa is His or Lys or Arg

<220>
<221> VARIANT
<222> (3)
<223> Wherein Xaa is His or Lys or Arg

<220>
<221> VARIANT
<222> (5)
<223> Wherein Xaa is not Pro

<220>
<221> VARIANT
<222> (6)
<223> Wherein Xaa is not Pro

<220>
<221> VARIANT
<222> (7)
<223> Wherein Xaa is Phe or Tyr or Trp

<220>
<221> VARIANT
<222> (8)
<223> Wherein Xaa is Phe or Tyr or Trp

<220>
<223> Description of Artificial Sequence: Consensus
sequence

<400> 12
Xaa Pro Xaa Pro Xaa Xaa Xaa
1 5

<210> 13
<211> 11
<212> PRT
<213> Artificial Sequence

<220>

<221> VARIANT

<222> (11)

<223> Wherein Xaa is not Met

<220>

<223> Description of Artificial Sequence:Consensus
sequence

<400> 13

Arg Pro Lys Pro Gln Gln Phe Phe Gly Leu Xaa

1

5

10

<210> 14

<211> 13

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)

<223> Wherein Xaa is Pyr or Tyr

<400> 14

Xaa Leu Tyr Glu Asn Lys Pro Arg Arg Pro Tyr Ile Leu

1

5

10

090907-14001